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Kevin R. Carman  
*Louisiana State University*

Jay C. Means  
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# Response of sedimentary bacteria in a Louisiana salt marsh to contamination by diesel fuel

Kevin R. Carman<sup>1,\*</sup>, Jay C. Means<sup>2</sup>, Steven C. Pomarico<sup>1</sup>

<sup>1</sup>Department of Zoology & Physiology, <sup>2</sup>Veterinary Physiology, Pharmacology & Toxicology, Louisiana State University, Baton Rouge, Louisiana 70803-1725, USA

**ABSTRACT:** In a 28 d microcosm study, we examined the effects of diesel-contaminated sediment on the sedimentary bacterial community of a Louisiana (USA) salt marsh that has been chronically exposed to petroleum hydrocarbons for decades. Diesel contaminants in microcosms as determined from polycyclic aromatic hydrocarbon (PAH) concentration ranged from 0.55 to 55 ppm (dry weight). Bacterial metabolism (incorporation of <sup>14</sup>C-acetate and <sup>3</sup>H-leucine) and bacterial abundance were not affected by diesel-contaminated sediment at any concentration. Bacterial degradation of <sup>14</sup>C-phenanthrene, however, increased in direct proportion to the amount of diesel-contaminated sediment added. Ambient sediment also exhibited significant capacity to degrade PAH. The half life of phenanthrene (based on <sup>14</sup>C-phenanthrene-degradation experiments) ranged from 137 d in ambient sediments to 4.5 d in sediment chronically exposed to high levels of diesel-contaminated sediments for 28 d. Two- and three-ring PAH, including naphthalenes, phenanthrenes, and dibenzothiophenes, constituted the bulk of PAH composition of diesel and were rapidly metabolized. Alkylated PAH were also readily metabolized. The rapid removal of PAH suggests that even if the marsh were exposed to chronically high levels of petroleum hydrocarbons, chemical evidence of the contaminants would not be detected in sediments. Collectively, these results are consistent with the hypothesis that the bacterial community in this salt marsh has adapted to chronic exposure to petroleum hydrocarbons.

**KEY WORDS:** Bacteria · Sediments · PAH · Diesel · Petroleum

## INTRODUCTION

It is estimated that  $1.7$  to  $8.8 \times 10^6$  t of petroleum hydrocarbons are released into the marine environment annually; 10% or more of this input may be from refined petroleum such as fuel oils (National Research Council 1985a). Among the various refined petroleum products, diesel fuel is considered to be highly toxic because it is enriched with polycyclic aromatic hydrocarbons (PAH; approximately 30 to 40%; National Toxicology Program 1986), the most toxic component of petroleum hydrocarbons (Clark 1989, Kennish 1992). Because of its toxicity and widespread use in military, commercial, and recreational vessels, diesel fuel represents a potentially significant contaminant to aquatic environments. Most of the PAH released into aquatic environments (approximately  $1.7 \times 10^5$  t yr<sup>-1</sup>) accumu-

lates in estuaries (Kennish 1992). As opposed to lighter fuels such as gasoline, many of the PAH in diesel are of a sufficiently high molecular weight that they do not readily evaporate (Clark 1989), but become associated with fine hydrophobic particles and are ultimately transported to the benthos (Connell & Miller 1984). Salt marshes are low-energy environments where these particles are likely to accumulate (Little 1987). Salt marshes are also highly productive and serve as nursery grounds for many commercially and economically important species. Because of these physical and biological characteristics, salt marshes are considered to be particularly susceptible to chronic and/or catastrophic inputs of petroleum hydrocarbons (National Research Council 1985b, Samiullah 1985).

Several studies have examined the response of benthic microbial communities to individual PAH (Bauer & Capone 1985a, b, Bauer et al. 1988, MacGillivray & Shiaris 1994) or crude oil (Alexander & Schwarz 1980,

\*E-mail: zocarm@lsuvm.sncc.lsu.edu

Griffiths et al. 1981a, b, Heitkamp & Cerniglia 1988), but few have considered the effects of refined fuels (Jamison et al. 1976). Further, while it is recognized that microorganisms play a critical role in the breakdown of hydrocarbons, the impact of hydrocarbons on the metabolism and abundance of natural microbial communities is poorly understood (Bartha & Atlas 1987). Crude oil, for example, has been shown to enhance (Bunch 1987), reduce (Griffiths et al. 1981a), or have no effect (Bauer & Capone 1985a, Wyndham 1985) on total abundance of sedimentary bacteria. Studies of individual aromatic compounds typically detect no significant influence on total bacterial abundance (Bauer & Capone 1985a). Bacterial communities also vary considerably in their metabolic response to petroleum hydrocarbons (Alexander & Schwarz 1980, Griffiths et al. 1981a, b, Bauer & Capone 1985a, Bauer et al. 1988). Previous chronic exposure to hydrocarbons has been proposed as a partial explanation for variability in bacterial response to petroleum hydrocarbons (Griffiths et al. 1981b).

This report is part of a study in which microcosm experiments were performed to examine the effects of diesel fuel on the benthic food web of a coastal salt marsh. Future papers will consider the impact of diesel on microalgal activity and abundance, meiofaunal grazing, and meiofaunal community structure. Here, we examine the influence of diesel-contaminated sediments on the benthic bacterial assemblage in terms of abundance, metabolic activity, and capacity to degrade PAH.

## MATERIALS AND METHODS

**Study site.** The research was performed using sediments from Terrebonne Bay estuary (29° 15' N, 91° 21' W) near the Louisiana Universities Marine Consortium Laboratory (LUMCON) at Cocodrie, LA, USA. Tidal range in the estuary is approximately 0.3 m and salinity ranges from 4 to 26 ppt. The estuary is a highly productive salt marsh that is dominated by the cord grass *Spartina alterniflora*. Sediment has a median grain size of 38 µm and is composed primarily of silts (41%) and clays (17%) (Chandler & Fleeger 1983). Organic content of sediment is approximately 2.5%. The study site is located in a region of intense hydrocarbon production and drilling activity, and commercial and recreational boat traffic is high. These combined factors lead to a high probability that the marsh experiences chronic exposure to both refined and crude hydrocarbons.

**Experimental design.** The effects of diesel fuel on sedimentary bacteria were examined using intact, natural sediment collected in cylindrical microcosms from

the study site. Microcosms were maintained in the LUMCON laboratory under controlled temperature and light conditions. Experimental treatments consisted of the daily addition to microcosms of small doses of diesel-contaminated sediment, and bacterial responses were determined over a 28 d period.

Microcosm experiments were performed with a 2 × 4 × 5 factorial design, with 2 wet tables (as blocks), 4 exposure times, and 5 diesel treatments as factors. Each diesel × time combination was replicated twice in both wet tables. Microcosms were constructed of 15.2 cm i.d. PVC pipe with windows covered with Nitex mesh (62 µm) to allow exchange of water. At low tide on 22 May 1994, 80 microcosms of exposed unvegetated sediment were collected by hand from mud flats surrounded by *Spartina alterniflora* marsh. Microcosms were gently pushed into the sediment to a depth of 15 cm, mud was excavated from the outside of the microcosm, and a form-fitting base was placed on the bottom. Intact microcosms were removed from the mud flat and transported to the LUMCON facility. Forty microcosms were randomly assigned to each of the 2 wet tables.

Microcosms were irrigated individually using a drip system. Ambient marsh water was filtered (5 µm) and pumped into a 1200 l holding tank. Water was aerated by continuous recirculation. Water was pumped from the holding tank to a 60 l head tank, which fed the drip system. Water was dripped into microcosms at a rate of approximately 1 l h<sup>-1</sup>, sufficient to exchange the overlying water approximately once every hour.

The treatments consisted of the addition to microcosms of sediment spiked with 3 levels of diesel (High, Medium, and Low), and 2 types of controls; in one control (Cont1), no sediment was added to microcosms, in the second control (Cont2) 'uncontaminated' sediment was added to microcosms. Four replicate microcosms (2 from each wet table) of each of the 5 treatment levels (20 total microcosms) were harvested at each of 4 time intervals (0, 7, 14, and 28 d) following a previously determined randomization schedule.

**Diesel-contaminated sediments.** Surficial sediments (top 2 cm) were collected from the marsh and processed following the procedure of Chandler (1986), which results in sterile sediment consisting of particles <62 µm. Diesel fuel was obtained from a commercial vendor. Two liters of processed sediments and 600 ml of diesel were placed in an amber 4 l bottle and tumbled for 10 d. The bottle was then removed from the tumbler and sediment allowed to settle overnight. Diesel was aspirated from the bottle and 1 l of 15 ppt artificial seawater (ASW) was added. The mixture was tumbled again (overnight), allowed to settle, and the supernatant aspirated. This procedure was repeated 3 times (total of 4 rinses). The sediment-water slurry

was transferred to 35 ml glass centrifuge tubes and centrifuged at  $1700 \times g$  for 3 min. The supernatant was removed and replaced with fresh ASW. Sediment and water were mixed thoroughly then recentrifuged. The supernatant was decanted again, and the process was repeated for a total of 4 rinses via centrifugation. Sediment was then recombined into a single batch and mixed to assure homogeneity. A sediment sample was removed from the batch, and total PAH (described below) was determined to be 687 ppm (dry weight). Contaminated sediment was then diluted with ambient sediment (processed as described above) to achieve PAH concentrations of 550, 55, and 5.5 ppm (dry weight). Diluted contaminated sediments were added to microcosms as described below with the objective of achieving final added concentrations in the top 1 cm of sediment of 55 (High), 5.5 (Medium), and 0.55 (Low) ppm.

At the beginning of the experiment, microcosms were dosed by adding sediment sufficient to create a 1 mm thick layer of sediment on the microcosm surface. This was accomplished by loading 30 ml plastic syringes with 17.8 ml of contaminated (or control) sediment, then slowly dispensing the sediment into the water overlying the microcosm (after removal of drip tubes) in a uniform manner. Sediment settled onto the microcosm surface within approximately 1 h, at which time microcosm drip tubes were replaced. Within approximately 2 h, surface topography (tubes, burrows, and tracks) from resident meiofauna and macrofauna was apparent. On each subsequent day, microcosms were dosed with 1.8 ml of sediment, sufficient to create a 0.1 mm sediment layer on the surface of microcosms.

Total PAH in sediment used to dose High treatments, as well as sediment in the top 1 cm of Day 0 and Day 28 High and Medium treatments, were determined with an Iatroscan (Ackman et al. 1990). For Iatroscan analysis, 10 to 34 g of sediment was extracted thrice in 70 ml dichloromethane, with 25 g of solvent-rinsed  $\text{Na}_2\text{SO}_4$  added in the first extraction.  $^{14}\text{C}$ -phenanthrene was added as an internal standard to determine extraction efficiency. Combined extracts were passed through a column containing 0.5 g solvent-rinsed  $\text{Na}_2\text{SO}_4$ , collected in a 250 ml round-bottom flask, and concentrated by rotary evaporation to 1–2 ml. The concentrated extract was transferred with rinsing to a  $13 \times 100$  mm tube then dried under  $\text{N}_2$  and stored in the freezer until further analysis. The extract was dissolved in  $\text{CHCl}_3$  and fractionated by solid phase extraction (SPE) chromatography on a silica column (500 mg, Whatman) with 5 ml  $\text{CHCl}_3$ . This fraction was dried under  $\text{N}_2$ , dissolved in toluene and fractionated by SPE chromatography on a silica column (500 mg) with 5 ml of toluene. The toluene fraction was dried

then dissolved in a 50–100  $\mu\text{l}$   $\text{CHCl}_3$  and duplicate 1–2  $\mu\text{l}$  samples were spotted on a Chromarod (SIII). Chromarods were dried under active vacuum after each development described below. Chromarod developments were carried out at  $35^\circ\text{C}$  as follows: (1) toluene for 5 min, (2) toluene for 5 min, (3) hexane for 30 min. Rods were analyzed using an Iatroscan MK-5 TLC/FID analyzer. The PAH peaks were quantified by comparison to an external calibration curve generated using a standard consisting of a mixture of 16 PAH ranging from naphthalene (2 rings) to benzo(g,h,i)perylene (6 rings) (Supelco). Final concentrations were calculated with a correction for recovery of  $^{14}\text{C}$ -phenanthrene.

Undiluted contaminated sediment and the top 1 cm of sediment from 2 replicates of each treatment on Day 0 and Day 28 were analyzed by gas chromatography/mass spectrometry (GC/MS) for PAH content (Means & McMillin 1993). Sediments were extracted in glass ointment jars containing ~4 g wet sediment after removing ~0.5 g for moisture determination.  $\text{Na}_2\text{SO}_4$  (30 g) was mixed into each sample and added to an empty container to create a reagent blank. Pesticide-grade dichloromethane (DCM, 40 ml) was added to each jar along with 15  $\mu\text{l}$  of a mixture of deuterated PAH (Ultra Scientific, Inc. #US-108) at  $40 \text{ ng } \mu\text{l}^{-1}$  in hexane. Open jars were placed in an ice-cooled sonicating bath for 12 min. Solvent was decanted through solvent-rinsed  $\text{Na}_2\text{SO}_4$  into a rotavap flask, and DCM extraction repeated twice more. Combined extracts were concentrated to ~1 ml, transferred with rinsing to a 4 ml vial, and further concentrated, with exchange to hexane, to 200  $\mu\text{l}$  using a dried nitrogen stream. Activated fine-granular copper (MacLeod et al. 1985) was added in excess to remove sulfur interference.

Extracts were analyzed by GC/MS using a Hewlett-Packard 5890/5970B Mass Selective Detector (MSD) equipped with a 30 m by 0.25 mm i.d., 0.25  $\mu\text{m}$  DB-5 film capillary column (J & W Scientific, Inc.). The GC was programmed from 50 to  $300^\circ\text{C}$  using 2 temperature ramps over a period of 60 min. The MSD was operated in selected-ion mode. Response calibration was achieved using a mixture of authentic reference standards at  $0.5 \text{ ng } \mu\text{l}^{-1}$ , which included parent PAH (US 106, Ultra Scientific), deuterated compounds (see above), and 42 alkylated naphthalenes, dibenzothiophenes, and phenanthrenes (Chiron Laboratories A.S., Norway). 2-Fluorobiphenyl was added to the extracts immediately prior to analysis to serve as an instrumental internal standard. Final concentrations were calculated with correction for recovery of the deuterated surrogate standards added during extraction.

**Direct counts.** Bacterial abundance in the top 1 cm of sediment was determined from acridine orange direct counts (AODC; Carman 1993). This procedure in-

cluded separation of bacteria from sediments by blending sediments in 0.01 % sodium pyrophosphate. The resulting supernatant was stained with 0.04 % acridine orange for 2 min, and bacteria were enumerated (Hobbie et al. 1977).

**<sup>14</sup>C-acetate incorporation.** Bacterial activity was measured by administering <sup>14</sup>C-acetate into sediment cores (1.7 cm i.d.) and following the label into bacterial membrane lipids (phospholipids) and lipid storage products (poly- $\beta$ -hydroxyalkanoates, PHA; Findlay & White 1987). Acetate was injected approximately 2 mm below the sediment-water interface through a silicon-sealed slit on the side of the core with a 50  $\mu$ l syringe (Hamilton; Dobbs et al. 1989). A 33.4 kBq quantity of [1,2-<sup>14</sup>C]acetate (dissolved in 22  $\mu$ l ASW; specific activity 4.0 GBq mmol<sup>-1</sup>) was added to each core and incubated in the dark (to prevent photosynthetic fixation of respired <sup>14</sup>CO<sub>2</sub>) for 5 h. Water overlying the sediment was discarded, and the top 1 cm of sediment was extruded into a glass 50 ml tube containing 25 ml of modified Bligh-Dyer Solution (White et al. 1979). Bulk lipids were extracted and then fractionated into neutral, phospho-, and glycolipids (which contain PHA)

(Guckert et al. 1985) and assayed for radioactivity. Controls were injected with <sup>14</sup>C-acetate and then immediately harvested as described above. Data were expressed as dpm <sup>14</sup>C incorporated after correction for controls.

**<sup>3</sup>H-leucine incorporation.** Cores of sediment (1.7 cm i.d.) were collected from microcosms on Day 28 only and injected with 405 kBq (22  $\mu$ l) of L-[4,5-<sup>3</sup>H]leucine (American Radiolabeled Chemicals, Inc.; 2.2 GBq mmol<sup>-1</sup>) as described above for <sup>14</sup>C-acetate incubations. Sediments were incubated for 45 min, after which the water overlying the sediment was discarded and the top 1 cm of sediment was extruded into a plastic bag. Sediment was then frozen and stored in liquid nitrogen until further processing. Controls were injected with <sup>3</sup>H-leucine and immediately frozen. Incorporation of <sup>3</sup>H into protein was determined after an acid/base hydrolysis procedure to separate protein from other macromolecules (Carman et al. 1988). Data were expressed as dpm <sup>3</sup>H incorporated after correction for controls.

**PAH metabolism.** Bacterial metabolism of PAH was examined with a modified version of the procedure

Table 1. Concentrations of parent and alkylated PAH as determined by GC/MS analysis. Diesel: sediment that was contaminated with diesel and added to microcosms in various dilutions (see text for further details). Control: the average of both types of control microcosms (2 samples each from Cont1 and Cont2). Low, Medium, and High: the 3 diesel treatments. Values are ppm (dry weight) and are the average of 2 replicates. Nap = naphthalene; Phen = phenanthrene; DBT = dibenzothiophene. % alkylated = proportion of total PAH that contained 1 or more alkyl side chains

Compound	Diesel	Control		Low		Medium		High	
		Day 0	Day 28	Day 0	Day 28	Day 0	Day 28	Day 0	Day 28
Nap	2601	3	4	0	2	12	0	2	8
C1-Nap	44054	2	2	0	2	90	0	47	5
C2-Nap	143537	2	1	0	1	395	0	186	8
C3-Nap	141640	7	6	0	10	1412	0	944	132
C4-Nap	84171	7	5	0	8	3717	0	3274	609
Fluorene	6843	0	3	1	0	8	0	2	1
Phen	17321	7	9	19	2	54	18	39	614
C1-Phen	53680	5	8	41	3	209	49	280	581
C2-Phen	56659	5	10	85	3	420	68	1314	1189
C3-Phen	38957	6	13	95	5	346	83	1210	1535
DBT	5169	0	1	0	0	12	0	17	26
C1-DBT	23930	1	52	2	0	105	0	155	210
C2-DBT	52212	6	12	46	6	427	15	1693	1175
Fluoranthene	825	49	25	37	18	87	61	41	46
Pyrene	3839	52	26	93	29	97	97	169	212
Benzantracene	105	11	98	19	34	37	43	11	34
Chrysene	26	39	32	52	12	55	22	31	15
Benzo(b)fluoranthene	7	41	29	45	44	66	59	25	18
Benzo(k)fluoranthene	2	5	20	30	40	26	39	2	12
Benzo(a)pyrene	5	13	14	21	13	34	37	6	12
Total parent	48548	228	287	323	210	549	374	349	1000
Total alkylated	638841	36	67	269	38	7119	214	9101	5442
% alkylated	92.9	13.5	18.8	45.5	15.3	92.8	36.4	96.3	84.5
Total PAH	687389	264	354	592	248	7668	587	9450	6442

described by MacGillivray & Shiaris (1994). Microcosm sediment was sampled with a 3 cc syringe core, and the top 1 cm (1 cm<sup>3</sup>) was extruded into sterile, 35 ml serum bottles. Nine ml of 0.45 µm filtered marsh water was added to bottles. Nine ml of 2% formaldehyde was added to controls. [9-<sup>14</sup>C]phenanthrene (Sigma, 307 MBq mmol<sup>-1</sup>) was solubilized in ethanol, and 10 µl (26.3 kBq; 0.085 µmol) was added to serum bottles. Serum bottles were capped with a rubber stopper through which a small plastic cup containing fluted Whatman #1 filter paper was inserted. Bottles were placed on a shaker table and incubated in the dark at 27°C for 72 h. Phenethylamine (0.1 ml) was injected into the wick, and 1.0 ml of 1 N HCl was added. Acidified samples were incubated overnight, after which wicks were removed and assayed for radioactivity. Data were expressed as percent <sup>14</sup>C-phenanthrene converted to CO<sub>2</sub> after subtraction of control values. Radioactivity recovered in control wicks averaged 1.3% (range 0.6 to 2.5%) of the total radioactivity added. Radioactivity in controls ranged from an average of 51% of radioactivity in all experimental values on Day 0 to <4% of radioactivity in High treatments on Days 14 and 28.

## RESULTS

### PAH composition

Absolute concentrations of major PAH classes are summarized in Table 1. The most abundant classes of PAH in diesel-contaminated sediment were naphthalenes, phenanthrenes, and dibenzothiophenes (DBT). Alkylated PAH made up 93% of the total PAH. The high proportions of naphthalenes, phenanthrenes, DBT and alkylated PAH are typical of refined petroleum hydrocarbons (National Research Council 1985c). To examine compositional differences in PAH among treatments, proportional PAH abundances were calculated (Steinhauer & Boehm 1992). The concentration of each compound was expressed as a fraction of the compound with the highest concentration. For example, in diesel-contaminated sediment used to dose microcosms, C2-naphthalenes had the highest concentration (Table 1) and all other PAH were expressed as a fraction of C2-naphthalene concentration (Fig. 1). In comparison to diesel-contaminated sediment, ambient sediment was relatively depleted in 2- and 3-ring PAH, and most PAH was in the form of 4- and 5-ring compounds (Fig. 1).

Composition of PAH in Day 0 microcosms was variable, but generally reflected that of the added diesel-contaminated sediment (Table 1, Fig. 2). In High (Fig. 2C) and Medium (Fig. 2B) treatments, naphtha-

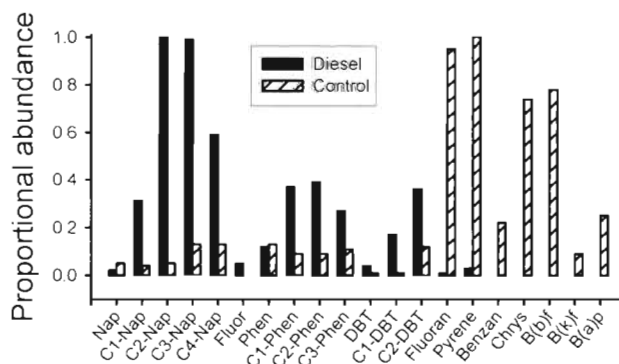


Fig. 1. Proportional abundance of major groups of PAH in diesel-contaminated and control (ambient) sediment. Nap = naphthalene, Fluor = fluorene, Phen = phenanthrene, DBT = dibenzothiophene, Fluoran = fluoranthene, Benzan = benzan-thracene, Chrys = chrysene, B(b)F = benzo(b)fluoranthene, B(k)F = benzo(k)fluoranthene, B(a)P = benzo(a)pyrene. C1, C2, C3, and C4: alkylated homologs of parent compounds containing from 1 to 4 methyl side chains, respectively. Values are averages of 2 replicates

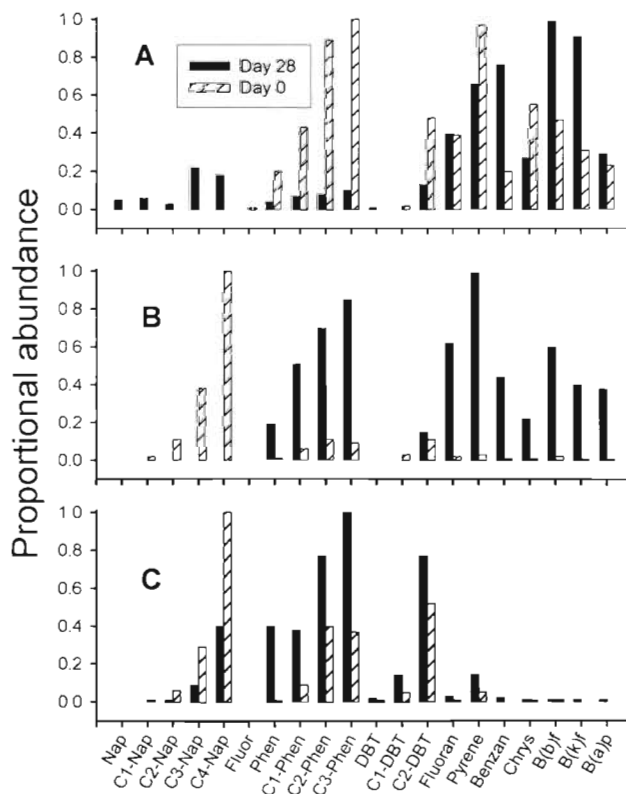


Fig. 2 Change in proportional abundance of major groups of PAH in (A) Low, (B) Medium, and (C) High treatments over the 28 d study period. Abbreviations as in Fig. 1. Values are averages of 2 replicates



lene and C1-, C2-, and C3-naphthalenes were proportionately less abundant than they were in diesel-contaminated sediments (Fig. 1). The loss of these lower-molecular-weight compounds in microcosms was probably the result of rapid sediment-water exchange and volatilization. Low treatments (Fig. 2A) consisted of only a minor PAH addition to microcosms, and thus the proportional abundances of PAH were similar to those of controls, i.e. very little naphthalene (parent or alkylated) and relatively high abundances of 4- and 5-ring compounds. The relatively high concentrations of phenanthrenes and C2-DBT were, however, evidence of the addition of diesel-contaminated sediments.

PAH composition in Day 28 microcosms differed substantially from Day 0 samples, and the degree of change differed among treatments (Fig. 2). In Day 28 Low treatments, the proportional abundance of phenanthrenes and C2-DBT was greatly reduced relative to Day 0 Low samples (Fig. 2A), and the profile closely resembled Day 0 control sediments (Fig. 1).

The alkylated naphthalenes that were abundant in Day 0 Medium treatments were completely eliminated by Day 28 (Fig. 2B). The proportional abundances of phenanthrenes and 4- and 5-ring compounds increased substantially by Day 28.

In High treatments, the compositional change of PAH from Day 0 to Day 28 was less dramatic than in Medium and Low treatments (Fig. 2C). Naphthalenes were reduced, but not eliminated as in Medium treatments. Phenanthrenes and DBT replaced naphthalenes as the dominant compounds, and, with the possible exception of pyrene, 4- and 5-ring PAH remained a minor component of total PAH composition.

### PAH concentration

Average total PAH (from GC/MS measurements) in control microcosms (Cont1 and Cont2) was 0.26 ppm on Day 0 and 0.35 ppm on Day 28. Total PAH in Day 0 Low, Medium, and High treatments were 0.59, 7.7, and 9.4 ppm respectively (Table 1). Concentrations in Low and Medium treatments were similar to expected concentrations (calculated PAH concentration of contaminated sediment plus ambient concentration), but the value in High treatments did not represent the 10× increase over Medium treatments that was expected. As discussed below, this apparent discrepancy may have resulted from uneven distribution of diesel-contaminated sediments in Day 0 microcosms.

In Medium treatments, concentration of all PAH containing ≤3 rings decreased from Day 0 to Day 28, whereas the concentration of 4/6 4- and 5-ring compounds increased (Table 1). With a few notable excep-

tions, changes in PAH concentrations in High treatments were similar to those observed in Medium treatments (Table 1). The concentrations of all alkylated naphthalenes decreased and, with the exception of C2-alkylated forms, the concentrations of all phenanthrenes and DBT decreased. As in Medium treatments 4/6 4- and 5-ring PAH increased in abundance over time in High treatments.

The concentration of 3-ring compounds decreased in Low treatments from Day 0 to Day 28 (Table 1). In contrast to Medium and High treatments, however, the concentrations of 4- and 5-ring compounds in Low treatments did not change substantially over time.

Measurements of total PAH by Iatroscan were generally higher than those determined from GC/MS, especially in the Day 0 High treatment. There are various possible explanations for this discrepancy. The first is related to calibration. The Iatroscan method that we used separated all PAH into a single peak. The PAH composition of our standard did not precisely match that of samples (the composition of which was variable), and variable detector responses to different compounds could have resulted in an overestimation of total PAH. Nevertheless, the measured total PAH concentration in Day 0 High treatments as determined from Iatroscan analysis (69.7 ppm) was reasonably close to the calculated expected concentration of 55 ppm. The apparent discrepancy between Iatroscan and GC/MS could also have been the result of the high degree of variability that was detected in Day 0 High treatments. For example, Iatroscan values from individual replicates of Day 0 High treatments were 169.2, 97.9, 6.0, and 5.8 ppm. This variability could have been the result of an uneven distribution of diesel-contaminated sediments in microcosms immediately after contaminated sediments were added to microcosms. The 2 GC/MS samples corresponded to the 6.0 and 169.2 ppm Iatroscan samples, and yielded values of 4.9 and 14.0 ppm respectively. Variability in Day 0 Iatroscan data from the High treatment could also have been a consequence of a loss of naphthalenes during processing of samples. We have observed that non-alkylated naphthalene is lost during the process of Chromarod development. Since naphthalene was less abundant after Day 0, this source of variability was probably less important after Day 0. Collectively, however, PAH as determined by GC/MS was a good predictor of total PAH as determined by Iatroscan ( $r^2 = 0.89$ , data not shown). Thus, at a minimum, the Iatroscan provided a good relative indication of total PAH concentration.

Iatroscan data indicated that PAH in High treatments accumulated over the first week, then decreased by approximately one-half by Day 14, and again by one-half by Day 28 (Fig. 3). Total PAH in Medium

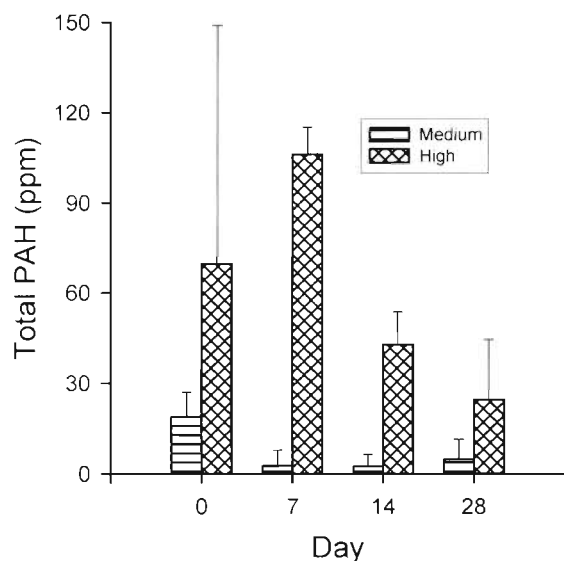


Fig. 3. Total PAH concentration in High and Medium treatments as determined by Iatroscan. Values are means + 1 SD (n = 4)

treatments decreased by a factor of approximately 7 by Day 7 and remained relatively constant thereafter (Fig. 3). These trends were qualitatively consistent with GC/MS data, which indicated that total PAH in Day 28 Medium treatments was only slightly greater than Day 0 controls, while total PAH in Day 28 High treatments was higher than Day 0 controls by a factor of 10 or more (Table 1). Thus, removal rate of PAH in Medium (and Low) treatments was equal to or exceeded the rate of addition. The removal rate of PAH from High treatments, however, was not sufficient to reduce PAH concentrations to background levels.

#### Bacterial abundance

Bacterial abundance in microcosms ranged from  $0.27$  to  $2.8 \times 10^9$  cells  $g^{-1}$  dry wt throughout the experiment (Fig. 4). Bacterial abundance was not significantly affected by diesel-contaminated sediment ( $p = 0.178$ ), and there was no trend that was even suggestive of an effect. Bacterial abundance did vary significantly among days ( $p < 0.0001$ ), with greatest numbers being detected on Day 7.

#### Bacterial activity

As with bacterial abundance, bacterial activity as determined by  $^{14}C$ -acetate incorporation into phospholipids, or the phospholipid:PHA (poly- $\beta$ -hydroxyalkanoates) ratio of  $^{14}C$ -acetate incorporation (Findlay & White 1987) were not significantly influenced by

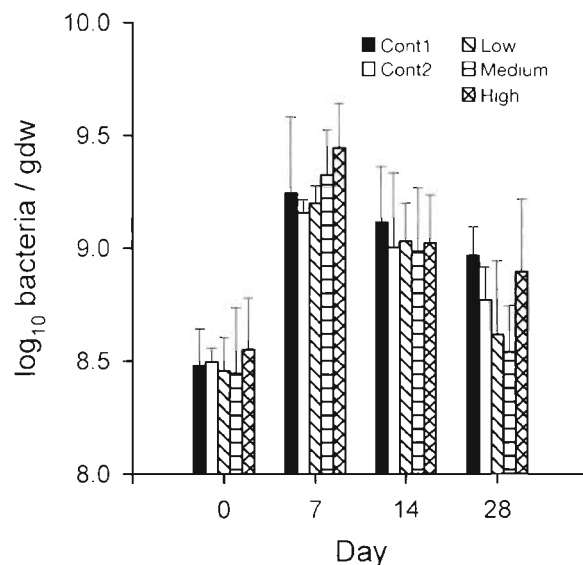


Fig. 4. Bacterial abundance in sediments exposed to a range of diesel contamination over a 28 d period. Values are means + 1 SD (n = 4)

diesel-contaminated sediment ( $p = 0.674$  and  $0.739$ , respectively; Fig. 5). Similarly,  $^3H$ -leucine incorporation into protein (measured on Day 28 only) was remarkably consistent among treatments (Fig. 6;  $p = 0.742$ ).

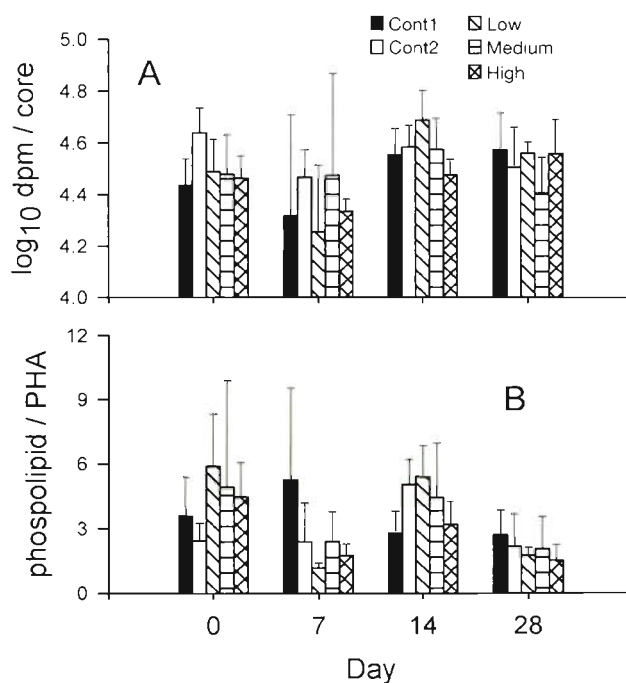


Fig. 5.  $^{14}C$ -acetate metabolism in sediments exposed to a range of diesel contamination over a 28 d period. (A) Incorporation of  $^{14}C$  into phospholipids. (B) Phospholipid/PHA ratio of  $^{14}C$  incorporation. Values are means + 1 SD (n = 4)



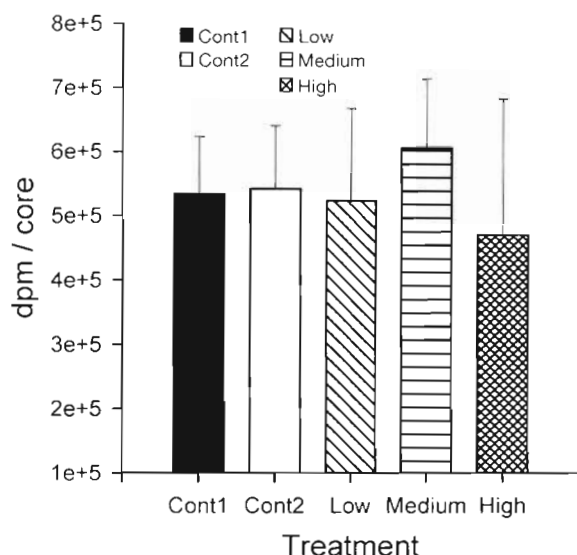


Fig. 6. <sup>3</sup>H-leucine incorporation into protein in sediments exposed to a range of diesel contamination on Day 28 of the experiment. Values are means + 1 SD (n = 4)

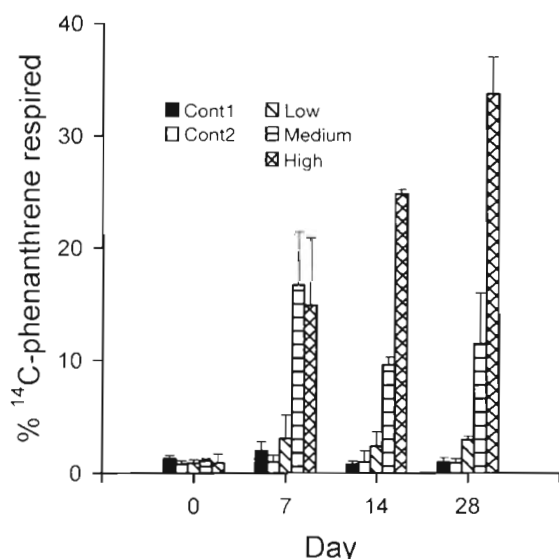


Fig. 7. Conversion of <sup>14</sup>C-phenanthrene to <sup>14</sup>CO<sub>2</sub> in sediments exposed to a range of diesel contamination over a 28 d period. Sediments were incubated for 72 h at 27°C. Values are means + 1 SD (n = 4)

#### PAH metabolism

In contrast to bacterial abundance and assays of bacterial activity, bacterial degradation of <sup>14</sup>C-phenanthrene was sensitive to diesel-contaminated sediment (Fig. 7). In Day 0 microcosms, degradation of <sup>14</sup>C-phenanthrene was low but detectable (0.9 to 1.3% of total available over a 72 h period) and did not differ among treatments. Degradation rates of <sup>14</sup>C-phenan-

threne remained relatively low in both controls (Cont1 and Cont2) over the entire course of the experiment (range 0.8 to 2.0%). Dose-dependent enhancement of phenanthrene degradation in all diesel treatments (Low, Medium, and High) occurred from Day 7 through Day 28. The enhancement of <sup>14</sup>C-phenanthrene degradation was statistically significant in Medium and High treatments when performing ANOVA on the entire data set, and when considering Days 7 through 28 individually (p < 0.0001). Degradation of <sup>14</sup>C-phenanthrene in Low treatments was significantly higher than in controls only on Day 28. After Day 7, the enhancement of <sup>14</sup>C-phenanthrene degradation in Low and Medium treatments remained constant or was slightly diminished. <sup>14</sup>C-phenanthrene degradation in High treatments continued to increase throughout the experimental period.

#### DISCUSSION

Our observations on the effects of diesel on Gulf of Mexico sedimentary bacteria appear to be generally consistent with previous studies of individual PAH or crude oils. Even at the highest doses (ca 55 ppm PAH), diesel-contaminated sediment had no detectable influence on bacterial incorporation of <sup>14</sup>C-acetate or <sup>3</sup>H-leucine, or on bacterial abundance. In a microcosm study similar in design to that reported here, PAH-contaminated sediments from a produced-water (contaminated water released from oil-production activities) site in the Gulf of Mexico also failed to elicit a change in sedimentary bacterial abundance or metabolism of <sup>14</sup>C-acetate (Carman et al. 1995). Nor did diesel-contaminated sediments have an influence on the physiological condition of the bacterial community as indicated by the relative incorporation of <sup>14</sup>C-acetate into phospholipids and PHA. Our experimental manipulations could have potentially produced both a physical (addition of sediment) and a chemical (addition of hydrocarbons) disturbance to benthic microorganisms. Nevertheless, no evidence of disturbance was detected.

Failure to detect changes in bacterial abundance or metabolic activity, however, does not mean that the bacterial community was unaffected by addition of hydrocarbons. Indeed, Baker & Griffiths (1984) proposed that evolved resistance to environmental contaminants may be responsible for variability in responses of sedimentary microorganisms to petroleum hydrocarbons. Further, Griffiths et al. (1981b) proposed that the insensitivity of Gulf of Mexico bacteria to petroleum hydrocarbons is the result of adaptation to chronic exposure from years of oil-production activities in the area.

Cerniglia & Heitkamp (1989) proposed that microbial adaptation to PAH contamination occurs as a 2-

step process: (1) acutely toxic low-molecular-weight PAH (such as naphthalene) eliminates sensitive microbes, and (2) resistant microbes that can metabolize PAH undergo a period of increased growth/activity. Our observations are partially consistent with this hypothesis. First, we observed no evidence of acute (or chronic) toxicity. Neither bacterial metabolic activity nor bacterial abundance was significantly influenced by diesel after short (Day 0 or Day 7) periods of exposure. Thus, any mortality or suppression of activity that might have occurred was below the sensitivity of our techniques. It is worth noting, however, that with approximately  $10^9$  bacteria  $\text{g}^{-1}$  of sediment, it is possible to eliminate or add millions of bacteria and not detect the change in a typical direct-count procedure. The second part of Cerniglia & Heitkamp's hypothesis is generally supported by our observations. Bacterial metabolism of  $^{14}\text{C}$ -phenanthrene dramatically increased over time in a dose-dependent pattern, implying that a PAH-degrading assemblage of bacteria developed in response to the presence of diesel-contaminated sediments. Again, however, we detected no significant change in total bacterial abundance that was related to diesel contamination, suggesting that (1) growth of PAH-degrading bacteria was offset by mortality of other bacteria, (2) existing bacteria have the capacity to metabolically switch to PAH degradation, or (3) the total number of PAH-degrading bacteria was insignificant relative to the total bacterial community.

Although  $^{14}\text{C}$ -phenanthrene degradation rates were relatively low at Day 0, degradation was nevertheless detectable ( $0.30$  to  $0.43\%$   $\text{d}^{-1}$ ). Using estuarine sediments from New York, Bauer & Capone (1985b) observed that  $^{14}\text{C}$ -anthracene (another 3-ring aromatic) degradation was only  $0.01\%$   $\text{d}^{-1}$  even after 4 d of exposure at 100 ppm. Thus, it would appear that ambient bacteria in this Louisiana salt marsh exhibit some significant level of preadaptation to PAH. Nevertheless, even modest ( $0.55$  ppm) additions of diesel elicited significant elevations of phenanthrene degradation. In the Low and Medium treatments, the rate of  $^{14}\text{C}$ -phenanthrene degradation peaked by 1 wk, and remained constant thereafter.  $^{14}\text{C}$ -phenanthrene degradation in High treatments, however, continued to increase throughout the experiment, and reached a rate of  $11.2\%$   $\text{d}^{-1}$  by Day 28. For comparison, Bauer & Capone (1985b) reported a maximum degradation rate for anthracene of  $3.9\%$   $\text{d}^{-1}$ . Further, the maximum phenanthrene degradation rates reported here are comparable to the maximum rates of naphthalene (a much more labile PAH) degradation ( $10\%$   $\text{d}^{-1}$ ) reported by Bauer & Capone (1985b).

The continued acceleration of phenanthrene degradation in High treatments apparently occurred because the supply of hydrocarbons outpaced the rate

at which they were metabolized. Specifically, PAH in Day 28 High treatments were still highly enriched in alkylated PAH, including naphthalenes, phenanthrenes, and DBT, indicating the presence of unmetabolized petroleum hydrocarbons.

We also observed that alkylated PAH, which are generally diagnostic of petroleum hydrocarbons, were readily removed from sediments. Relatively little information is available concerning the metabolism of alkylated versus parent PAH. Cerniglia & Heitkamp (1989) observed that 2-methylnaphthalene was metabolized much slower than naphthalene or phenanthrene. Our data suggest that even highly methylated naphthalene (C4) was readily metabolized, as were alkylated forms of other PAH (i.e. phenanthrenes and dibenzothiophenes). In Low and Medium treatments, parent and alkylated naphthalenes, phenanthrenes, and DBT were removed completely or almost completely over the 28 d study period. The rate of decrease in parent phenanthrene and DBT in Low and Medium treatments was lower than the rates of decrease in alkylated forms. In the High treatment, parent phenanthrene and DBT increased by approximately a factor of 15 and 1.4 respectively, whereas accumulation of alkylated phenanthrenes and DBT was generally much lower (Table 1). This implies that the removal of PAH was not simply a desorption phenomenon (Means et al. 1980, Means & Wijayarathne 1984, Means & McMillin 1993, Means in press); if such were the case, higher molecular weight (alkylated) compounds would have been removed more slowly. Thus, microbial degradation must have contributed significantly to the removal of 2- and 3-ring parent and alkylated PAH, and alkylated PAH showed no evidence of being disproportionately resistant to microbial degradation.

Previous studies have suggested that DBT may provide a reliable marker for petroleum-hydrocarbon contamination because they are found in all types of petroleum (Clark 1989, Steinhauer et al. 1994), including diesel (Williams et al. 1986), and they are considered to be resistant to photochemical (Andersson 1993) and microbial (Sinkkonen 1989) degradation. In the present study, however, DBT were at least as susceptible to microbial degradation as were phenanthrenes. Over the 28 d study, essentially all DBT were removed from Low and Medium treatments, thus leaving no evidence of diesel contamination. Fayad & Overton (1995) also observed high rates of DBT degradation in sediments contaminated during the 1991 Gulf War. In particular, they observed that C2- and C3-DBT were degraded more quickly than C1-DBT, an observation that is consistent with our data (Table 1). Our data also show that, in High treatments, the rate of accumulation of DBT was much less than that of phenanthrenes. Thus, our data indicate that DBT were metabolized by sedimen-

tary bacteria at a high rate, and that DBT would not accumulate in these sediments unless very high rates of input were maintained.

Collectively, our data indicate that the Louisiana salt marsh bacterial community studied here is symptomatic of one that has been chronically exposed to petroleum hydrocarbons: bacterial abundance and general assays of bacterial metabolism are insensitive to additions of diesel, ambient bacteria can metabolize PAH at substantial rates, and the PAH-degrading portion of the community responds quickly to additions of petroleum hydrocarbons. It is possible that the PAH-degrading bacterial community maintains ambient sedimentary PAH concentrations at relatively low levels. In Low and Medium treatments, no significant accumulation of PAH could be detected over the 28 d period relative to ambient sediment. Further, PAH concentration in High treatments were reduced by approximately 43% over the 28 d experiment, despite the daily addition of diesel-contaminated sediment.

These observations of the bacterial response to diesel contamination have implications for understanding how ecosystems respond to contamination by crude or refined petroleum hydrocarbons. One possibility is that rapid bacterial metabolism of petroleum hydrocarbons could ultimately reduce exposure of other benthic organisms to potentially toxic compounds. We have observed that the meiofaunal/microbial foodweb in this salt marsh is relatively resistant to petroleum-hydrocarbon contamination from produced water (Carman et al. 1995) or diesel (Carman unpubl.). Further study will be required to determine if the apparent insensitivity of this benthic food web is because the fauna itself is resistant to hydrocarbons, or if it relies on bacterial detoxification of petroleum contaminants.

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